

GenCore version 4.5
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117: gb_est48:*
118: gb_est49:*
119: gb_est50:*
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122: gb_est53:*
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185: em.gss.hum4:*
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187: em.gss.inv2:*
188: em.gss.inv3:*
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232: gb.gss32:*
233: gb.gss33:*
234: gb.gss34:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
C 1	70.2	3.1	1101	CNS0039G	AJ063921	Drosophili
C 2	61.4	2.7	1101	CNS00EVL	AJ069706	Drosophili
C 3	61	2.7	1201	CNS0167M	AJJ06396	Drosophili
C 4	60.6	823	225	A2697380	A2687380	ENVLO56TP
C 5	60.2	2.6	1092	CNS020K7	AJJ175696	Tetraodon
C 6	60.2	2.6	1101	CNS01219	AJJ1595	Drosophili
C 7	60	2.6	1101	CNS0029	AJ069493	Drosophili
C 8	59.2	2.6	1032	CNS020IP	AJ206746	Tetraodon
C 9	58.6	2.6	1200	CNS016CO	AJJ06578	Drosophili
C 10	58	2.6	668	A254901	AJJ14901	Drosophili
C 11	57.8	2.5	891	A2693582	A2683582	ENTRK47TR
C 12	57.4	2.5	652	CNS03HOU	AJ244551	Tetraodon
C 13	57.4	2.5	1101	CNS008B	AJ063632	Drosophili
C 14	56.8	2.5	1101	CNS00KA	AJ077628	Drosophili
C 15	56.4	2.5	928	BF274633	BF274633	GA_Eb002
C 16	56.2	2.5	881	A2547573	A2547573	ENTGB68IR
C 17	56	2.5	1101	CNS00B07	AJ069440	Drosophili
C 18	55.8	2.5	611	AA549883	AA549883	0956m3 gm

FEATURES	Location/Qualifiers
SOURCE	1928

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Source          /organism="Gossypium arboreum"
                /strain="ATA"
                /cultivar="8400"
                /db_xref="taxon:29729"
                /clone="GA_Eb0021J19f"
                /clone.lib="Gossypium arboreum 7-10 dpa fiber library"
                /tissue_type="fibers isolated from bolls harvested 7-10
dpa"
                /lab.host="E. coli"
                /note="vector: PBK-CMV; Site_1: EcoRI; Site_2: XbaI"
                /origin="a" 85 c 80 g 248 t 15 others
                /base_count=500

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Search completed: June 6, 2001, 20:03:01
Job time: 4842 sec